RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/537,520_

Source: _PCT/10_

Date Processed by STIC: _6/14/05_

ENTERED

Raw Sequence Listing before editing,
for reference only

PCT

RAW SEQUENCE LISTING   DATE: 06/14/2005
PATENT APPLICATION: US/10/537,520   TIME: 09:05:45

Input Set : A:\3125USOP_SEQ.txt
Output Set: N:\CRF4\06142005\J537520.raw

```
          2 <110> APPLICANT: KAJINO, Masahiro
          3        NAKAYAMA, Yutaka
          4        KIMURA, Haruhide
          6 <120> TITLE OF INVENTION: 1,3-Benzothiazinone derivatives, production and use thereof
          8 <130> FILE REFERENCE: 3125 USOP
C-->     10 <140> CURRENT APPLICATION NUMBER: US/10/537,520
C-->     10 <141> CURRENT FILING DATE: 2005-06-03
         10 <150> PRIOR APPLICATION NUMBER: PCT/JP2003/015535
         11 <151> PRIOR FILING DATE: 2003-12-04
         13 <150> PRIOR APPLICATION NUMBER: JP2002-353546
         14 <151> PRIOR FILING DATE: 2002-12-05
W-->     16 <150> PRIOR APPLICATION NO: PCT/JP2003/015535
         17 <151> PRIOR FILING DATE: 2003-12-04
         19 <160> NUMBER OF SEQ ID NOS: 2
```

delete – already shown above

ERRORED SEQUENCES

Does Not Comply
Corrected Diskette Needed

```
         29 <210> SEQ ID NO: 2
         30 <211> LENGTH: 42
         31 <212> TYPE: DNA
         32 <213> ORGANISM: Artificial Sequence
         34 <220> FEATURE:
         35 <223> OTHER INFORMATION: mutant ARE
         37 <400> SEQUENCE: 2
         38 gagcttggaa atggcattgc taatggtggc aaagcaactt tg          42
E-->     40 1/1
E-->     45 1
```

VERIFICATION SUMMARY   DATE: 06/14/2005
PATENT APPLICATION: **US/10/537,520**   TIME: 09:05:46

Input Set : **A:\3125USOP_SEQ.txt**
Output Set: **N:\CRF4\06142005\J537520.raw**

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:16 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:40 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:43 SEQ:2
L:40 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
M:254 Repeated in SeqNo=2
L:45 M:252 E: No. of Seq. differs, <211> LENGTH:Input:42 Found:43 SEQ:2